



#7

SEQUENCE LISTING

<110> OWMAN, CHRISTER

<120> HEPTAHELIX RECEPTOR AND ITS USE AS LEUKOTRIENE B4  
RECEPTOR

<130> 07675.0001-03 SEQUENCE LISTING

<140> 09/893,512

<141> 2001-06-29

<150> 60/061,789

<151> 1997-10-14

<150> 60/081,958

<151> 1998-04-15

<150> 09/170,069

<151> 1998-10-13

<160> 17

<170> PatentIn Ver. 2.1

<210> 1

<211> 1672

<212> DNA

<213> Homo sapiens

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 ggcgggagtg gagtggaaga agagggagag gtggagcaaa gtgagggccg agtgagagcg 1620  
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 <212> PRT  
 <213> Homo sapiens

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*Sub D7*

Ile Ser Leu Leu Ala Ile Ile Leu Leu Ser Val Ala Leu Ala Val Gly  
 20 25 30

Leu Pro Gly Asn Ser Phe Val Val Trp Ser Ile Leu Lys Arg Met Gln  
 35 40 45

Lys Arg Ser Val Thr Ala Leu Met Val Leu Asn Leu Ala Leu Ala Asp  
 50 55 60

Leu Ala Val Leu Leu Thr Ala Pro Phe Phe Leu His Phe Leu Ala Gln  
 65 70 75 80

Gly Thr Trp Ser Phe Gly Leu Ala Gly Cys Arg Leu Cys His Tyr Val  
 85 90 95

Cys Gly Val Ser Met Tyr Ala Ser Val Leu Leu Ile Thr Ala Met Ser  
 100 105 110

Leu Asp Arg Ser Leu Ala Val Ala Arg Pro Phe Val Ser Gln Lys Leu  
 115 120 125

Arg Thr Lys Ala Met Ala Arg Arg Val Leu Ala Gly Ile Trp Val Leu  
 130 135 140

Ser Phe Leu Leu Ala Thr Pro Val Leu Ala Tyr Arg Thr Val Val Pro  
 145 150 155 160

Trp Lys Thr Asn Met Ser Leu Cys Phe Pro Arg Tyr Pro Ser Glu Gly  
165 170 175

His Arg Ala Phe His Leu Ile Phe Glu Ala Val Thr Gly Phe Leu Leu  
180 185 190

Pro Phe Leu Ala Val Val Ala Ser Tyr Ser Asp Ile Gly Arg Arg Leu  
195 200 205

Gln Ala Arg Arg Phe Arg Arg Ser Arg Arg Thr Gly Arg Leu Val Val  
210 215 220

Leu Ile Ile Leu Thr Phe Ala Ala Phe Trp Leu Pro Tyr His Val Val  
225 230 235 240

Asn Leu Ala Glu Ala Arg Arg Ala Leu Ala Gly Gln Ala Ala Gly Leu  
245 250 255

Gly Leu Val Gly Lys Arg Leu Ser Leu Ala Arg Asn Val Leu Ile Ala  
260 265 270

Leu Ala Phe Leu Ser Ser Ser Val Asn Pro Val Leu Tyr Ala Cys Ala  
275 280 285

Gly Gly Gly Leu Leu Arg Ser Ala Gly Val Gly Phe Val Ala Lys Leu  
290 295 300

Leu Glu Gly Thr Gly Ser Glu Ala Ser Ser Thr Arg Arg Gly Gly Ser  
305 310 315 320

Leu Gly Gln Thr Ala Arg Ser Gly Pro Ala Ala Leu Glu Pro Gly Pro  
325 330 335

Ser Glu Ser Leu Thr Ala Ser Ser Pro Leu Lys Leu Asn Glu Leu Asn  
340 345 350

<210> 3

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<212> DNA

<213> Homo sapiens

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<210> 4  
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<210> 5  
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<210> 6  
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<211> 8  
<212> PRT  
<213> Homo sapiens

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1 5

<210> 8  
<211> 18  
<212> PRT  
<213> Homo sapiens

<220>  
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<223> Xaa at position 6 is any amino acid

<220>  
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<222> (7)  
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<222> (17)  
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1 5 10 15

Xaa Trp

<210> 9  
<211> 350  
<212> PRT  
<213> Homo sapiens

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Met Ser Asn Ile Thr Asp Pro Gln Met Trp Asp Phe Asp Asp Leu Asn  
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Phe Thr Gly Met Pro Pro Ala Asp Glu Asp Tyr Ser Pro Cys Met Leu  
20 25 30

Glu Thr Glu Thr Leu Asn Lys Tyr Val Val Ile Ile Ala Tyr Ala Leu  
35 40 45

Val Phe Leu Leu Ser Leu Leu Gly Asn Ser Leu Val Met Leu Val Ile  
50 55 60

Leu Tyr Ser Arg Val Gly Arg Ser Val Thr Asp Val Tyr Leu Leu Asn  
65 70 75 80

Leu Ala Leu Ala Asp Leu Leu Phe Ala Leu Thr Leu Pro Ile Trp Ala  
85 90 95

Ala Ser Lys Val Asn Gly Trp Ile Phe Gly Thr Phe Leu Cys Lys Val  
100 105 110

Val Ser Leu Leu Lys Glu Val Asn Phe Tyr Ser Gly Ile Leu Leu Leu  
115 120 125

Ala Cys Ile Ser Val Asp Arg Tyr Leu Ala Ile Val His Ala Thr Arg  
130 135 140

Thr Leu Thr Gln Lys Arg His Leu Val Lys Phe Val Cys Leu Gly Cys  
145 150 155 160

Trp Gly Leu Ser Met Asn Leu Ser Leu Pro Phe Phe Leu Phe Arg Gln  
165 170 175

*Sub D7* Ala Tyr His Pro Asn Asn Ser Ser Pro Val Cys Tyr Glu Val Leu Gly  
180 185 190

Asn Asp Thr Ala Lys Trp Arg Met Val Leu Arg Ile Leu Pro His Thr  
195 200 205

Phe Gly Phe Ile Val Pro Leu Phe Val Met Leu Phe Cys Tyr Gly Phe  
210 215 220

Thr Leu Arg Thr Leu Phe Lys Ala His Met Gly Gln Lys His Arg Ala  
225 230 235 240

Met Arg Val Ile Phe Ala Val Val Leu Ile Phe Leu Leu Cys Trp Leu  
245 250 255

Pro Tyr Asn Leu Val Leu Leu Ala Asp Thr Leu Met Arg Thr Gln Val  
260 265 270

Ile Gln Glu Thr Cys Glu Arg Arg Asn Asn Ile Gly Arg Ala Leu Asp  
275 280 285

Ala Thr Glu Ile Leu Gly Phe Leu His Ser Cys Leu Asn Pro Ile Ile  
290 295 300

Tyr Ala Phe Ile Gly Gln Asn Phe Arg His Gly Phe Leu Lys Ile Leu  
305 310 315 320

Ala Met His Gly Leu Val Ser Lys Glu Phe Leu Ala Arg His Arg Val  
325 330 335

Thr Ser Tyr Thr Ser Ser Ser Val Asn Val Ser Ser Asn Leu  
340 345 350

<210> 10

<211> 355

<212> PRT

<213> Homo sapiens

<400> 10

Met Glu Ser Asp Ser Phe Glu Asp Phe Trp Lys Gly Glu Asp Leu Ser  
1 5 10 15

Asn Tyr Ser Tyr Ser Ser Thr Leu Pro Pro Phe Leu Leu Asp Ala Ala  
20 25 30

Pro Cys Glu Pro Glu Ser Leu Glu Ile Asn Lys Tyr Phe Val Val Ile  
35 40 45

Ile Tyr Ala Leu Val Phe Leu Leu Ser Leu Leu Gly Asn Ser Leu Val  
50 55 60

Met Leu Val Ile Leu Tyr Ser Arg Val Gly Arg Ser Val Thr Asp Val  
65 70 75 80

Tyr Leu Leu Asn Leu Ala Leu Ala Asp Leu Leu Phe Ala Leu Thr Leu  
85 90 95

Pro Ile Trp Ala Ala Ser Lys Val Asn Gly Trp Ile Phe Gly Thr Phe  
100 105 110

Leu Cys Lys Val Val Ser Leu Leu Lys Glu Val Asn Phe Tyr Ser Gly  
115 120 125

Ile Leu Leu Leu Ala Cys Ile Ser Val Asp Arg Tyr Leu Ala Ile Val  
130 135 140

His Ala Thr Arg Thr Leu Thr Gln Lys Arg Tyr Leu Val Lys Phe Ile  
145 150 155 160

Cys Leu Ser Ile Trp Gly Leu Ser Leu Leu Leu Ala Leu Pro Val Leu

165								170				175			
Leu	Phe	Arg	Arg	Thr	Val	Tyr	Ser	Ser	Asn	Val	Ser	Pro	Ala	Cys	Tyr
180								185				190			
Glu	Asp	Met	Gly	Asn	Asn	Thr	Ala	Asn	Trp	Arg	Met	Leu	Leu	Arg	Ile
195								200				205			
Leu	Pro	Gln	Ser	Phe	Gly	Phe	Ile	Val	Pro	Leu	Leu	Ile	Met	Leu	Phe
210								215				220			
Cys	Tyr	Gly	Phe	Thr	Leu	Arg	Thr	Leu	Phe	Lys	Ala	His	Met	Gly	Gln
225								230				235			
Lys	His	Arg	Ala	Met	Arg	Val	Ile	Phe	Ala	Val	Val	Leu	Ile	Phe	Leu
245								250				255			
Leu	Cys	Trp	Leu	Pro	Tyr	Asn	Leu	Val	Leu	Leu	Ala	Asp	Thr	Leu	Met
260								265				270			
Arg	Thr	Gln	Val	Ile	Gln	Glu	Thr	Cys	Glu	Arg	Arg	Asn	His	Ile	Asp
275								280				285			
Arg	Ala	Leu	Asp	Ala	Thr	Glu	Ile	Leu	Gly	Ile	Leu	His	Ser	Cys	Leu
290								295				300			
Asn	Pro	Leu	Ile	Tyr	Ala	Phe	Ile	Gly	Gln	Lys	Phe	Arg	His	Gly	Leu
305								310				315			
Leu	Lys	Ile	Leu	Ala	Ile	His	Gly	Leu	Ile	Ser	Lys	Asp	Ser	Leu	Pro
325								330				335			
Lys	Asp	Ser	Arg	Pro	Ser	Phe	Val	Gly	Ser	Ser	Ser	Gly	His	Thr	Ser
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Thr	Thr	Leu													
355															

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 Met Glu Thr Pro Asn Thr Thr Glu Asp Tyr Asp Thr Thr Thr Glu Phe  
 1 5 10 15





Cys Glu Gln Ser Arg His Leu Asp Leu Ala Val Gln Val Thr Glu Val  
275 280 285

Ile Ala Tyr Thr His Cys Cys Val Asn Pro Val Ile Tyr Ala Phe Val  
290 295 300

Gly Glu Arg Phe Arg Lys Tyr Leu Arg Gln Leu Phe His Arg Arg Val  
305 310 315 320

Ala Val His Leu Val Lys Trp Leu Pro Phe Leu Ser Val Asp Arg Leu  
325 330 335

Glu Arg Val Ser Ser Thr Ser Pro Ser Thr Gly Glu His Glu Leu Ser  
340 345 350

Ala Gly Phe  
355

<210> 12

<211> 379

<212> PRT

<213> Homo sapiens

<400> 12

Pro Glu Pro Met Glu Thr Pro Asn Thr Thr Glu Asp Tyr Asp Thr Thr  
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Thr Glu Phe Asp Tyr Gly Asp Ala Thr Pro Cys Gln Lys Val Asn Glu  
20 25 30

Arg Ala Phe Gly Ala Gln Leu Leu Pro Pro Leu Tyr Ser Leu Val Phe  
35 40 45

Val Ile Gly Leu Val Pro Glu Pro Gly Asn Ile Leu Val Val Leu Val  
50 55 60

Leu Val Gln Tyr Lys Arg Leu Lys Asn Met Thr Ser Ile Tyr Leu Leu  
65 70 75 80

Asn Leu Ala Ile Ser Asp Leu Leu Phe Leu Phe Thr Leu Pro Phe Trp  
85 90 95

Ile Asp Tyr Lys Leu Lys Asp Asp Trp Val Pro Glu Pro Phe Gly Asp  
100 105 110

Ala Met Cys Lys Ile Leu Ser Gly Phe Tyr Tyr Thr Gly Leu Tyr Ser  
115 120 125

Glu	Ile	Phe	Phe	Ile	Ile	Leu	Leu	Thr	Ile	Asp	Arg	Tyr	Leu	Ala	Ile	
130						135					140					
Val	His	Ala	Val	Phe	Ala	Leu	Arg	Ala	Arg	Thr	Val	Thr	Phe	Gly	Pro	
145					150					155					160	
Glu	Pro	Val	Ile	Thr	Ser	Ile	Ile	Ile	Trp	Ala	Leu	Ala	Ile	Leu	Ala	
				165					170						175	
Ser	Met	Pro	Gly	Leu	Tyr	Phe	Ser	Lys	Thr	Gln	Trp	Glu	Phe	Thr	His	
			180					185					190			
His	Thr	Cys	Ser	Leu	His	Phe	Pro	His	Glu	Ser	Leu	Arg	Glu	Trp	Lys	
	195						200					205				
Leu	Phe	Gln	Ala	Pro	Glu	Pro	Leu	Lys	Leu	Asn	Leu	Phe	Gly	Leu	Val	
	210					215					220					
Leu	Pro	Leu	Leu	Val	Met	Ile	Ile	Cys	Tyr	Thr	Gly	Ile	Ile	Lys	Ile	
225					230					235					240	
Leu	Leu	Arg	Arg	Pro	Asn	Glu	Lys	Lys	Ser	Lys	Ala	Val	Arg	Leu	Ile	
				245					250					255		
Phe	Val	Ile	Met	Ile	Ile	Phe	Phe	Leu	Pro	Glu	Pro	Phe	Trp	Thr	Pro	
			260					265					270			
Tyr	Asn	Leu	Thr	Ile	Leu	Ile	Ser	Val	Phe	Gln	Asp	Phe	Leu	Phe	Thr	
	275						280					285				
His	Glu	Cys	Glu	Gln	Ser	Arg	His	Leu	Asp	Leu	Ala	Val	Gln	Val	Thr	
	290					295					300					
Glu	Val	Ile	Ala	Tyr	Thr	His	Cys	Cys	Val	Asn	Pro	Val	Ile	Pro	Glu	
305					310					315				320		
Pro	Tyr	Ala	Phe	Val	Gly	Glu	Arg	Phe	Arg	Lys	Tyr	Leu	Arg	Gln	Leu	
				325					330					335		
Phe	His	Arg	Arg	Val	Ala	Val	His	Leu	Val	Lys	Trp	Leu	Pro	Phe	Leu	
				340				345					350			
Ser	Val	Asp	Arg	Leu	Asp	Arg	Val	Ser	Ser	Thr	Ser	Pro	Ser	Thr	Gly	
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Glu	His	Glu	Pro	Glu	Pro	Leu	Ser	Ala	Gly	Phe						
370						375										

<210> 13  
<211> 374  
<212> PRT  
<213> Homo sapiens

<400> 13

Met Leu Ser Thr Ser Arg Ser Arg Phe Ile Arg Asn Thr Asn Glu Ser  
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Gly Glu Glu Val Thr Thr Phe Phe Asp Tyr Asp Tyr Gly Ala Pro Cys  
20 25 30

His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu Pro Pro Leu  
35 40 45

Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn Met Leu Val Val  
50 55 60

Leu Ile Leu Ile Asn Cys Lys Lys Leu Lys Cys Leu Thr Asp Ile Tyr  
65 70 75 80

Leu Leu Asn Leu Ala Ile Ser Asp Leu Leu Phe Leu Ile Thr Leu Pro  
85 90 95

Leu Trp Ala His Ser Ala Ala Asn Glu Trp Val Phe Gly Asn Ala Met  
100 105 110

Cys Lys Leu Phe Thr Gly Leu Tyr His Ile Gly Tyr Phe Gly Gly Ile  
115 120 125

Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu Ala Ile Val His  
130 135 140

Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe Gly Val Val Thr  
145 150 155 160

Ser Val Ile Thr Trp Leu Val Ala Val Phe Ala Ser Val Pro Gly Ile  
165 170 175

Ile Phe Thr Lys Cys Gln Lys Glu Asp Ser Val Tyr Val Cys Gly Pro  
180 185 190

Tyr Phe Pro Arg Gly Trp Asn Asn Phe His Thr Ile Met Arg Asn Ile  
195 200 205

Leu Gly Leu Val Leu Pro Leu Leu Ile Met Val Ile Cys Tyr Ser Gly

210	215	220
Ile Leu Lys Thr Leu Leu Arg Cys Arg Asn Glu Lys Lys Arg His Arg		
225	230	235 240
Ala Val Arg Val Ile Phe Thr Ile Met Ile Val Tyr Phe Leu Phe Trp		
	245	250 255
Thr Pro Tyr Asn Ile Val Ile Leu Leu Asn Thr Phe Gln Glu Phe Phe		
	260	265 270
Gly Leu Ser Asn Cys Glu Ser Thr Ser Gln Leu Asp Gln Ala Thr Gln		
	275	280 285
Val Thr Glu Thr Leu Gly Met Thr His Cys Cys Ile Asn Pro Ile Ile		
	290	295 300
Tyr Ala Phe Val Gly Glu Lys Phe Arg Ser Leu Phe His Ile Ala Leu		
	305	310 315 320
Gly Cys Arg Ile Ala Pro Leu Gln Lys Pro Val Cys Gly Gly Pro Gly		
	325	330 335
Val Arg Pro Gly Lys Asn Val Lys Val Thr Thr Gln Gly Leu Leu Asp		
	340	345 350
Gly Arg Gly Lys Gly Lys Ser Ile Gly Arg Ala Pro Glu Ala Ser Leu		
	355	360 365
Gln Asp Lys Glu Gly Ala		
	370	

<210> 14  
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 <213> Homo sapiens

<400> 14  
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 20 25 30  
 Ala Pro Cys His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu  
 35 40 45

Pro	Pro	Leu	Tyr	Ser	Pro	Glu	Pro	Leu	Val	Phe	Ile	Phe	Gly	Phe	Val
50						55					60				
Gly	Asn	Met	Leu	Val	Val	Leu	Ile	Leu	Ile	Asn	Cys	Lys	Lys	Leu	Lys
65					70					75					80
Cys	Leu	Thr	Asp	Ile	Tyr	Leu	Leu	Asn	Leu	Ala	Ile	Ser	Asp	Leu	Leu
				85					90					95	
Phe	Leu	Ile	Thr	Leu	Pro	Leu	Trp	Ala	His	Pro	Glu	Pro	Ser	Ala	Ala
			100					105					110		
Asn	Glu	Trp	Val	Phe	Gly	Asn	Ala	Met	Cys	Lys	Leu	Phe	Thr	Gly	Leu
			115				120					125			
Tyr	His	Ile	Gly	Tyr	Phe	Gly	Gly	Ile	Phe	Phe	Ile	Ile	Leu	Leu	Thr
	130					135					140				
Ile	Asp	Arg	Tyr	Leu	Ala	Ile	Val	His	Ala	Val	Phe	Ala	Leu	Lys	Pro
145					150					155					160
Glu	Pro	Ala	Arg	Thr	Val	Thr	Phe	Gly	Val	Val	Thr	Ser	Val	Ile	Thr
				165					170					175	
Trp	Leu	Val	Ala	Val	Phe	Ala	Ser	Val	Pro	Gly	Ile	Ile	Phe	Thr	Lys
			180					185					190		
Cys	Gln	Lys	Glu	Asp	Ser	Val	Tyr	Val	Cys	Gly	Pro	Tyr	Phe	Pro	Arg
		195					200					205			
Gly	Trp	Asn	Asn	Pro	Glu	Pro	Phe	His	Thr	Ile	Met	Arg	Asn	Ile	Leu
	210					215					220				
Gly	Leu	Val	Leu	Pro	Leu	Leu	Ile	Met	Val	Ile	Cys	Tyr	Ser	Gly	Ile
225					230					235					240
Leu	Lys	Thr	Leu	Leu	Arg	Cys	Arg	Asn	Glu	Lys	Lys	Arg	His	Arg	Ala
				245					250					255	
Val	Arg	Val	Ile	Phe	Thr	Ile	Met	Ile	Pro	Glu	Pro	Val	Tyr	Phe	Leu
			260					265					270		
Phe	Trp	Thr	Pro	Tyr	Asn	Ile	Val	Ile	Leu	Leu	Asn	Thr	Phe	Gln	Glu
	275					280						285			
Phe	Phe	Gly	Leu	Ser	Asn	Cys	Glu	Ser	Thr	Ser	Gln	Leu	Asp	Gln	Ala
	290					295					300				

Put D7

Thr Gln Val Thr Glu Thr Leu Gly Met Thr His Cys Cys Ile Pro Glu  
305 310 315 320

Pro Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe Arg Arg Tyr  
325 330 335

Leu Ser Val Phe Phe Arg Lys His Ile Thr Lys Arg Phe Cys Lys Gln  
340 345 350

Cys Pro Val Phe Tyr Arg Glu Thr Val Asp Gly Val Thr Ser Thr Asn  
355 360 365

Thr Pro Ser Pro Glu Pro Thr Gly Glu Gln Glu Val Ser Ala Gly Leu  
370 375 380

<210> 15

<211> 350

<212> PRT

<213> Homo sapiens

<400> 15

Met Asn Ser Phe Asn Tyr Thr Thr Pro Asp Tyr Gly His Tyr Asp Asp  
1 5 10 15

Lys Asp Thr Leu Asp Leu Asn Thr Pro Val Asp Lys Thr Ser Asn Thr  
20 25 30

Leu Arg Val Pro Asp Ile Leu Ala Leu Val Ile Phe Ala Val Val Phe  
35 40 45

Leu Val Gly Val Leu Gly Asn Ala Leu Val Val Trp Val Thr Ala Phe  
50 55 60

Glu Ala Lys Arg Thr Ile Asn Ala Ile Trp Phe Leu Asn Leu Ala Val  
65 70 75 80

Ala Asp Phe Leu Ser Cys Leu Ala Leu Pro Ile Leu Phe Thr Ser Ile  
85 90 95

Val Gln His His His Trp Pro Phe Gly Gly Ala Ala Cys Ser Ile Leu  
100 105 110

Pro Ser Leu Ile Leu Leu Asn Met Tyr Ala Ser Ile Leu Leu Leu Ala  
115 120 125

15

Thr Ile Ser Ala Asp Arg Phe Leu Leu Val Phe Lys Pro Ile Trp Cys  
130 135 140

Gln Asn Phe Arg Gly Ala Gly Leu Ala Trp Ile Ala Cys Ala Val Ala  
145 150 155 160

Trp Gly Leu Ala Leu Leu Leu Thr Ile Pro Ser Phe Leu Tyr Arg Val  
165 170 175

Val Arg Glu Glu Tyr Phe Pro Pro Lys Val Leu Cys Gly Val Asp Tyr  
180 185 190

Ser His Asp Lys Arg Arg Glu Arg Ala Val Ala Ile Val Arg Leu Val  
195 200 205

Leu Gly Phe Leu Trp Pro Leu Leu Thr Leu Thr Ile Cys Tyr Thr Phe  
210 215 220

Ile Leu Leu Arg Thr Trp Ser Arg Arg Ala Thr Arg Ser Thr Lys Thr  
225 230 235 240

Leu Lys Val Val Val Ala Val Val Ala Ser Phe Phe Ile Phe Trp Leu  
245 250 255

Pro Tyr Gln Val Thr Gly Ile Met Met Ser Phe Leu Glu Pro Ser Ser  
260 265 270

Pro Thr Phe Leu Leu Leu Asn Lys Leu Asp Ser Leu Cys Val Ser Phe  
275 280 285

Ala Tyr Ile Asn Cys Cys Ile Asn Pro Ile Ile Tyr Val Val Ala Gly  
290 295 300

Gln Gly Phe Gln Gly Arg Leu Arg Lys Ser Leu Pro Ser Leu Leu Arg  
305 310 315 320

Asn Val Leu Thr Glu Glu Ser Val Val Arg Glu Ser Lys Ser Phe Thr  
325 330 335

Arg Ser Thr Val Asp Thr Met Ala Gln Lys Thr Gln Ala Val  
340 345 350

<210> 16

<211> 351

<212> PRT

<213> Homo sapiens



<400> 16

Met Glu Thr Asn Ser Ser Leu Pro Thr Asn Ile Ser Gly Gly Thr Pro  
1 5 10 15

Ala Val Ser Ala Gly Tyr Leu Phe Leu Asp Ile Ile Thr Tyr Leu Val  
20 25 30

Phe Ala Val Thr Phe Val Leu Gly Val Leu Gly Asn Gly Leu Val Ile  
35 40 45

Trp Val Ala Gly Phe Arg Met Thr His Thr Val Thr Thr Ile Ser Tyr  
50 55 60

Leu Asn Leu Ala Val Ala Asp Phe Cys Phe Thr Ser Thr Leu Pro Phe  
65 70 75 80

Phe Met Val Arg Lys Ala Met Gly Gly His Trp Pro Phe Gly Trp Phe  
85 90 95

Leu Cys Lys Phe Leu Phe Thr Ile Val Asp Ile Asn Leu Phe Gly Ser  
100 105 110

Val Phe Leu Ile Ala Leu Ile Ala Leu Asp Arg Cys Val Cys Val Leu  
115 120 125

His Pro Val Trp Thr Gln Asn His Arg Thr Val Ser Leu Ala Lys Lys  
130 135 140

Val Ile Ile Gly Pro Trp Val Met Ala Leu Leu Leu Thr Leu Pro Val  
145 150 155 160

Ile Ile Arg Val Thr Thr Val Pro Gly Lys Thr Gly Thr Val Ala Cys  
165 170 175

Thr Phe Asn Phe Ser Pro Trp Thr Asn Asp Pro Lys Glu Arg Ile Asn  
180 185 190

Val Ala Val Ala Met Leu Thr Val Arg Gly Ile Ile Arg Phe Ile Ile  
195 200 205

Gly Phe Ser Ala Pro Met Ser Ile Val Ala Val Ser Tyr Gly Leu Ile  
210 215 220

Ala Thr Lys Ile His Lys Gln Gly Leu Ile Lys Ser Ser Arg Pro Leu  
225 230 235 240

Arg Val Leu Ser Phe Val Ala Ala Ala Phe Phe Leu Cys Trp Ser Pro

	245		250		255										
Tyr	Gln	Val	Val	Ala	Leu	Ile	Ala	Thr	Val	Arg	Ile	Arg	Glu	Leu	Leu
	260						265						270		
Gln	Gly	Met	Tyr	Lys	Glu	Ile	Gly	Ile	Ala	Val	Asp	Val	Thr	Ser	Ala
	275						280						285		
Leu	Ala	Phe	Phe	Asn	Ser	Cys	Leu	Asn	Pro	Met	Leu	Tyr	Val	Phe	Met
	290						295						300		
Gly	Gln	Asp	Phe	Arg	Glu	Arg	Leu	Ile	His	Ala	Leu	Pro	Ala	Ser	Leu
305					310					315				320	
Glu	Arg	Ala	Leu	Thr	Glu	Asp	Ser	Thr	Gln	Thr	Ser	Asp	Thr	Ala	Thr
				325					330					335	
Asn	Ser	Thr	Leu	Pro	Ser	Ala	Glu	Val	Ala	Leu	Gln	Ala	Lys	Cys	
			340					345					350		

<210> 17  
 <211> 352  
 <212> PRT  
 <213> Homo sapiens

<400> 17

Met	Asn	Thr	Thr	Ser	Ser	Ala	Ala	Pro	Pro	Ser	Leu	Gly	Val	Glu	Phe
1				5				10				15			
Ile	Ser	Leu	Leu	Ala	Ile	Ile	Leu	Leu	Ser	Val	Ala	Leu	Ala	Val	Gly
			20					25				30			
Leu	Pro	Gly	Asn	Ser	Phe	Val	Val	Trp	Ser	Ile	Leu	Lys	Arg	Met	Gln
		35				40					45				
Lys	Arg	Ser	Val	Thr	Ala	Leu	Met	Val	Leu	Asn	Leu	Ala	Leu	Ala	Asp
	50					55					60				
Leu	Ala	Val	Leu	Leu	Thr	Ala	Pro	Phe	Phe	Leu	His	Phe	Leu	Ala	Gln
65					70				75					80	
Gly	Thr	Trp	Ser	Phe	Gly	Leu	Ala	Gly	Cys	Arg	Leu	Cys	His	Tyr	Val
				85					90				95		
Cys	Gly	Val	Ser	Met	Tyr	Ala	Ser	Val	Leu	Leu	Ile	Thr	Ala	Met	Ser
				100				105					110		

Leu Asp Arg Ser Leu Ala Val	Ala Arg Pro Phe Val Ser Gln Lys Leu
115	120 125
Arg Thr Lys Ala Met Ala Arg	Arg Val Leu Ala Gly Ile Trp Val Leu
130	135 140
Ser Phe Leu Leu Ala Thr Pro Val	Leu Ala Tyr Arg Thr Val Val Pro
145	150 155 160
Trp Lys Thr Asn Met Ser Leu Cys	Phe Pro Arg Tyr Pro Ser Glu Gly
165	170 175
His Arg Ala Phe His Leu Ile Phe	Glu Ala Val Thr Gly Phe Leu Leu
180	185 190
Pro Phe Leu Ala Val Val Ala Ser	Tyr Ser Asp Ile Gly Arg Arg Leu
195	200 205
Gln Ala Arg Arg Phe Arg Arg Ser	Arg Arg Thr Gly Arg Leu Val Val
210	215 220
Leu Ile Ile Leu Thr Phe Ala Ala	Phe Trp Leu Pro Tyr His Val Val
225	230 235 240
Asn Leu Ala Glu Ala Arg Arg Ala	Leu Ala Gly Gln Ala Ala Gly Leu
245	250 255
Gly Leu Val Gly Lys Arg Leu Ser	Leu Ala Arg Asn Val Leu Ile Ala
260	265 270
Leu Ala Phe Leu Ser Ser Ser Val	Asn Pro Val Leu Tyr Ala Cys Ala
275	280 285
Gly Gly Gly Leu Leu Arg Ser Ala	Gly Val Gly Phe Val Ala Lys Leu
290	295 300
Leu Glu Gly Thr Gly Ser Glu Ala Ser	Ser Thr Arg Arg Gly Gly Ser
305	310 315 320
Leu Gly Gln Thr Ala Arg Ser Gly Pro	Ala Ala Leu Glu Pro Gly Pro
325	330 335
Ser Glu Ser Leu Thr Ala Ser Ser Pro	Leu Lys Leu Asn Glu Leu Asn
340	345 350

*End 77*